

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 09/765,555C  
Source: 1Fw/6  
Date Processed by STIC: 3/15/06

***ENTERED***



IFW16

## RAW SEQUENCE LISTING

DATE: 03/15/2006

PATENT APPLICATION: US/09/765,555C

TIME: 11:14:37

Input Set : E:\27801-20014.20 - Substitute Seqlist (2).txt  
 Output Set: N:\CRF4\03152006\I765555C.raw

3 <110> APPLICANT: BARBAS, Carlos  
 4 STEGE, Justin  
 5 GUAN, Xueni  
 6 DALMIA, Bipin  
 8 <120> TITLE OF INVENTION: METHODS AND COMPOSITIONS TO MODULATE  
 9 EXPRESSION IN PLANTS  
 11 <130> FILE REFERENCE: 27801-20014.20  
 13 <140> CURRENT APPLICATION NUMBER: 09/765,555C  
 14 <141> CURRENT FILING DATE: 2001-01-19  
 16 <150> PRIOR APPLICATION NUMBER: 09/620,897  
 17 <151> PRIOR FILING DATE: 2000-07-21  
 19 <150> PRIOR APPLICATION NUMBER: US 60/177,468  
 20 <151> PRIOR FILING DATE: 2000-01-21  
 22 <160> NUMBER OF SEQ ID NOS: 79  
 24 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
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 27 <211> LENGTH: 532  
 28 <212> TYPE: DNA  
 29 <213> ORGANISM: Artificial Sequence  
 31 <220> FEATURE:  
 32 <223> OTHER INFORMATION: Promoter CsVMV  
 34 <400> SEQUENCE: 1  
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 36 acggggaaaaa ctatggaagt attatgtgag ctcagcaaga agcagatcaa tatgcggcac 120  
 37 atatgcaacc tatgttcaaa aatgaagaat gtacagatac aagatcctat actgccagaa 180  
 38 tacgaagaag aatacgtaga aattgaaaaaa gaagaaccag gcgaagaaaa gaatcttgaa 240  
 39 gacgttaagca ctgacgacaa caatgaaaaa aagaagataa ggtcggtat tgtgaaagag 300  
 40 acatagagga cacatgtaag gtggaaaaatg taagggcgg aagtaacctt atcacaaagg 360  
 41 aatcttatcc cccactactt atccctttat attttccgt gtcattttg cccttgagtt 420  
 42 ttccatatata aggaaccaag ttccggcattt gtgaaaacaa gaaaaaaattt ggtgtaagct 480  
 43 attttctttg aagtactgag gatacaactt cagagaaatt tgtaagttt ta 532  
 45 <210> SEQ ID NO: 2  
 46 <211> LENGTH: 18  
 47 <212> TYPE: DNA  
 48 <213> ORGANISM: Artificial Sequence  
 50 <220> FEATURE:  
 51 <223> OTHER INFORMATION: Zinc finger protein 2C7 binding site  
 53 <400> SEQUENCE: 2  
 54 gcgtgggcgg cgtggcgg 18  
 56 <210> SEQ ID NO: 3  
 57 <211> LENGTH: 51  
 58 <212> TYPE: DNA  
 59 <213> ORGANISM: Artificial Sequence

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 62 <223> OTHER INFORMATION: Promoter pc7rbTATA  
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 68 <211> LENGTH: 3121  
 69 <212> TYPE: DNA  
 70 <213> ORGANISM: Artificial Sequence  
 72 <220> FEATURE:  
 73 <223> OTHER INFORMATION: pND3008 coding region  
 75 <400> SEQUENCE: 4  
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 77 aaatttaccac atatttttt tgcacactt gtttgaagtg cagtttatct atctttatac 120  
 78 atatattaa actttactct acgaataata taatctatag tactacaata atatcagtgt 180  
 79 ttttagagaat catataatg aacagttaga catggctaa aggacaattt agtattttga 240  
 80 caacaggact ctacagtttt atcttttttag tgcattgtt ttctcctttt tttttgcaaa 300  
 81 tagttcacc tatataatac ttcatccatt ttatttagtac atccattttttag gttttagggt 360  
 82 taatgggttt tatagactaa ttttttttagt acatctattt tattctattt tagctctaa 420  
 83 attaagaaaa ctaaaactct attttagttt ttttatttaa taatttagat attaaataga 480  
 84 attaaaataaa gtgactaaaa attaaacaaa tacccttaa gaaattaaaa aaactaaggaa 540  
 85 aacattttc ttgttcgag tagataatgc cagcctgtt aacgcgcgtcg acgagtctaa 600  
 86 cggacaccaa ccagcgaacc agcagcgtcg cgtcgggcca agcgaaggcag acggcacggc 660  
 87 atctctgtcg ctgcctctgg acccctctcg agagttccgc tccaccgttg gacttgctcc 720  
 88 gctgtcggca tccagaaaatt gctgtggcgga gcggcagacg tgagccggca cggcaggcgg 780  
 89 cctccctcctc ctctcacggc acggcagcta cgggggattt cttcccccacc gctcctcgc 840  
 90 tttcccttcc tcgccccccg taataaaatag acacccctc cacaccctct ttcccaacc 900  
 91 tcgtgttgcg cggagcgcac acacacacaa ccagatctcc cccaaatcca cccgtcggca 960  
 92 cctccgttcc aaggtacgccc gtcgtcctc ccccccccccc cctctctacc ttctctagat 1020  
 93 cggcggtccg gtccatgggtt agggcccggtt agttctactt ctgttcatgt ttgtgtttaga 1080  
 94 tccgtgtttt tgtagatcc gtgcgtctag cgttgcata cggatgcgac ctgtacgtca 1140  
 95 gacacgttct gattgctaact ttgcctgtt ttctctttgg gaaatctgg gatggctcta 1200  
 96 gccgttccgc agacgggatc gatttcatga tttttttgtt ttcgttgcattt agggtttgg 1260  
 97 ttgcccccttcc ctttatttcc aatataatgcc gtgcacttgtt ttgtcggttgc atcttttcat 1320  
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 101 tacatgttgc tgcgggtttt actgtatgc atacagagat gctttttgtt cgcttgggtt 1560  
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 104 agttacgagt ttaagatggaa tggaaatatc gatctaggat aggtatacat gttgtgtgg 1740  
 105 gtttactga tgcataataca tgcgttgcata tgcgttgcata attcatatgc tctaaccctt 1800  
 106 agtacctatc tattataata aacaagtatg ttttataattt attttgcattt tgatataactt 1860  
 107 ggatgtggc atatgcacgca gctatatgtt gatgttttgc gcccgttgc cttatcgat 1920  
 108 ttatgttgc tttttgttgc tgcgttgcata tgcgttgcata gttgttggt gttacttctg 1980  
 109 caggtcgact ctagaggatc tatggcccaag gcggccctcg agtcccttgc tgcgttgcata 2040  
 110 gtcgagtcctt gcgatcgccg cttttctaaag tgcgttgcata tgcgttgcata tgcgttgcata 2100  
 111 cacacaggcc agaaggccctt ccagtgtcga atatgcatgc gtaacttcag tgcgttgcata 2160  
 112 caccatccca cccacatccca caccacacaca ggcgagaagc cttttgccttgc tgcgttgcata 2220  
 113 gggaggaatg ttgcaggatc tgatgttgc aagaggatc cccaaatcca taccgggttgc 2280

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114	aagccctatg	cttgcctgt	cgagtccgc	gatgcgcgt	tttctaagtc	ggctgatctg	2340
115	aagcgccata	tccgcatcca	cacaggccag	aagcccttc	agtgtcgaaat	atgcatgcgt	2400
116	aacttcagtc	gtagtgcacca	ccttaccacc	cacatccgca	cccacacagg	cgagaagcct	2460
117	tttgcctgtg	acatttgtgg	gaggaagt	gccaggagtg	atgaacgcaa	gaggcatacc	2520
118	aaaatccatt	taagacagaa	ggactctaga	actagtggcc	aggccggca	ggctagcccg	2580
119	aaaaagaaaac	gcaaagttgg	gcgcgcccac	gchgctggacg	atttcgatct	cgacatgctg	2640
120	ggttctgatg	ccctcgatga	ctttgacctg	gatatgttgg	gaagcgacgc	attggatgac	2700
121	tttgatctgg	acatgctcg	ctccgatgct	ctggacgatt	tcgatctcg	tatgttaatt	2760
122	aactaccgt	acgacgttcc	ggactacgct	tcttgagaat	tcgcggccgc	ggggcccgagc	2820
123	ctagggagga	gctcaagatc	ccccgaattt	ccccgatcgt	tcaaacattt	ggcaataaaag	2880
124	tttcttaaga	ttgaatcctg	ttgcccgtct	tgcgatgatt	atcatctaat	ttctgttcaa	2940
125	ttacgttaag	catgtataaa	ttaacatgt	atgcgtacg	ttatttatga	gatgggtttt	3000
126	tatgattaga	gtcccgaat	tatacattt	atacgcgtat	gaaaacaaaa	tatagcgcgc	3060
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140	aaatttaccac	atattttttt	tgtcacactt	gtttaagt	cagtttatct	atctttatac	120
141	atatatattaa	actttactct	acgaataata	taatctatag	tactacaata	atatcagtgt	180
142	tttagagaat	catataaata	acagatgt	catgtctaa	aggacaattg	agtatatttga	240
143	caacaggact	ctacagttt	atcttttag	tgtgcgtgt	ttctccttt	tttttgc当地	300
144	tagttcacc	tatataatac	ttcatccatt	ttatttagtac	atccatttag	gtttagggt	360
145	taatggttt	tatagactaa	tttttttagt	acatctattt	tattctattt	tagctctaa	420
146	attaagaaaa	ctaaaactct	attttagttt	ttttattttaa	taatttagat	ataaaataga	480
147	ataaaataaa	gtgactaaaa	attaaacaaa	taccctttaa	gaaattaaaa	aaactaagga	540
148	aacatttttc	ttgtttcgag	tagataatgc	cagctgtt	aacgcccgt	acgagtctaa	600
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151	gctgtcggca	tccagaaatt	gcgtggcgga	gcggcagacg	tgagccgca	cgccaggcgg	780
152	cctcctcctc	ctctcacggc	acggcagcta	cgggggattt	cttcccacc	gctcccttcgc	840
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155	cctccgcttc	aaggtaacg	gccgtcctc	cccccccccc	cctctctacc	ttctcttagat	1020
156	cggcggtcc	gtccatgg	agggcccggt	agttctactt	ctgttcatgt	ttgtgtttaga	1080
157	tcgtgtttt	tgtagatcc	gtgtgtctag	cggtcgatca	cgatgcgtac	ctgtacgtca	1140
158	gacacgttct	gattgttaac	ttgtccagt	ttctttttgg	gaaatctgg	gatggctcta	1200
159	gccgtccgc	agacgggatc	gatttcatga	ttttttttgt	ttcgttgc	agggtttgg	1260
160	ttgccccttt	ccttatttc	aatatatgc	gtgcactt	ttgtcggtt	atctttcat	1320
161	gtttttttt	gtttttttt	tgatgtatgt	gtttttttgg	gcggcggtt	tagatcgag	1380
162	tagaattctg	tttcaacta	cctgggttgg	ttatattt	tggatctgt	tgtgtgtg	1440
163	atacatattc	atagttacga	attgaagatg	atggatggaa	atatcgatct	aggataggt	1500
164	tacatgttga	tgcgggtttt	actgtatgt	atacagatgt	gtttttgtt	cgcttgggtt	1560
165	tgtatgttga	gtgtgggttgg	gcggcggtt	attcgatct	gatcgagta	gaataactgtt	1620

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166	tcaaactacc	ttgtgttattt	attaattttg	gaactgtatg	tgtgtgtcat	acatcttcat	1680
167	agttacgagt	ttaagatgga	tggaaatatac	gatctaggat	aggatacat	gttcatgtgg	1740
168	gttttactga	tgcataataca	tgtatggcata	tgcagcatct	attcatatgc	tctaaccctt	1800
169	agtagctatc	tattataata	aacaagtatg	ttttataatt	attttgatct	tgatataactt	1860
170	ggatgatggc	atatgcagca	gctatatgtg	gatttttta	gccctgcctt	catacgctat	1920
171	ttatattgctt	ggtaactgttt	cttttgcga	tgctcaccct	gttgggggt	gttacttctg	1980
172	cagggtcgact	ctagaggatc	cactagttag	ccatgggcta	gcatggccgc	tgccgtgcgc	2040
173	atgaacatcc	agatgctgct	cgaagccgct	gattatctgg	aacgcccggga	gcgcgaagcc	2100
174	gagcacggct	acgcccagcat	gctgccatat	ccgaaaaaga	aacgcaaggt	ggcccaggcg	2160
175	gccctcgagc	tcccttatgc	ttgccctgtc	gagtcctgcg	atgcgcgtt	ttctaagtgc	2220
176	gtgtatctga	agcgcataat	ccgcataccac	acaggccaga	agcccttcca	gtgtcgaata	2280
177	tgcatgcgta	acttcagtcg	tagtgaccac	cttaccaccc	acatccgcac	ccacacaggc	2340
178	gagaagcctt	ttgcctgtga	catttgggg	aggaagttt	ccaggagtg	tgaacgcaag	2400
179	aggcataccca	aaatccatac	cggtgagaag	ccctatgtt	gccctgtcg	gtccctgcgat	2460
180	cggcgctttt	ctaagtcggc	tgtatctgaag	cgcataatcc	gcatccacac	agggcagaag	2520
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182	atccgcaccc	acacaggcga	gaagcctttt	gcctgtgaca	tttggggag	gaagtttgcc	2640
183	aggagtgtatg	aacgcgaaag	gcataccaaa	atccattaa	gacagaagga	ctctagaact	2700
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185	accgagctcg	gatccccca	atccccca	tcgttcaa	atttggcaat	aaagtttctt	2820
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187	taagcatgta	ataattaaca	tgtatgc	gacgttattt	atgagatgg	ttttatgtat	2940
188	tagagtcccg	caattataca	ttaataacgc	gatagaaaac	aaaatatacg	gcgcaaacta	3000
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195	<213>	ORGANISM:	Artificial Sequence				
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198	<223>	OTHER INFORMATION:	6X2C7 binding site				
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202	ggcggcgtgg	gcgactagtg	ctagcgcgt	ggcggcgtgg	gcaacaagc	gtggcgccg	120
203	tgggcgaaca	agcgtggcgc	gcgtggcga	ctagtg			156
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218	<212>	TYPE:	DNA				
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222	<223>	OTHER INFORMATION:	ZFP from -85 to -65				

RAW SEQUENCE LISTING DATE: 03/15/2006  
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232 <220> FEATURE:  
233 <223> OTHER INFORMATION: ZFPm1 from -68 to -85  
235 <400> SEQUENCE: 9  
236 tgagaggagg aaggaggc 18  
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239 <211> LENGTH: 18  
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243 <220> FEATURE:  
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247 gagtgagagg aggaagga 18  
249 <210> SEQ ID NO: 11  
250 <211> LENGTH: 24  
251 <212> TYPE: DNA  
252 <213> ORGANISM: Artificial Sequence  
254 <220> FEATURE:  
255 <223> OTHER INFORMATION: ZFP from 294 to 317  
257 <400> SEQUENCE: 11  
258 gccaactact acggctccct cacc 24  
260 <210> SEQ ID NO: 12  
261 <211> LENGTH: 18  
262 <212> TYPE: DNA  
263 <213> ORGANISM: Artificial Sequence  
265 <220> FEATURE:  
266 <223> OTHER INFORMATION: ZFPm3 from 311 to 294  
268 <400> SEQUENCE: 12  
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272 <211> LENGTH: 18  
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276 <220> FEATURE:  
277 <223> OTHER INFORMATION: ZFPm4 from 317 to 300  
279 <400> SEQUENCE: 13  
280 ggtgagggag ccgtagta 18  
282 <210> SEQ ID NO: 14  
283 <211> LENGTH: 3300  
284 <212> TYPE: DNA  
285 <213> ORGANISM: Artificial Sequence  
287 <220> FEATURE:  
288 <223> OTHER INFORMATION: Partial sequence of pMal-m1 and zinc finger  
289 protein ZFPm1

**VERIFICATION SUMMARY**

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